

Exhibit F

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<!--StartFragment-->RESULT 1
US-09-997-722-215
; Sequence 215, Application US/09997722
; Publication No. US20040072154A1
; GENERAL INFORMATION:
; APPLICANT: Morris, David
; APPLICANT: Engelhard, Eric
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
; FILE REFERENCE: A-71171/RMS/DCF
; CURRENT APPLICATION NUMBER: US/09/997,722
; CURRENT FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: US 09/747,377
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 301
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 215
; LENGTH: 2119
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-997-722-215

Query Match          100.0%; Score 2119; DB 3; Length 2119;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  GGCACGAGTAGGGGTGGCGGGTCAGTGTCTCGGGGGCTTCATCCAGGTCCCTGGA 60
Db      1  GGCACGAGTAGGGGTGGCGGGTCAGTGTCTCGGGGGCTTCATCCAGGTCCCTGGA 60

Qy     61  GTTCCTGGTCCCTGGAGCTCCGCAGTTGGCCGCCAACCTGGGTGAGGCAGCGGAGCTGTG 120
Db     61  GTTCCTGGTCCCTGGAGCTCCGCAGTTGGCCGCCAACCTGGGTGAGGCAGCGGAGCTGTG 120

Qy    121  GCGACTGGCCGGCCATGCTTCCCGGGCTGAGGACTATGAAGTGTGTACACCATTTGCA 180
Db    121  GCGACTGGCCGGCCATGCTTCCCGGGCTGAGGACTATGAAGTGTGTACACCATTTGCA 180

Qy    181  CAGGCTCCTACGGCCGCTGCCAGAAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTT 240
Db    181  CAGGCTCCTACGGCCGCTGCCAGAAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTT 240

Qy    241  GGAAGAACTTGACTATGGCTCCATGACAGAAAGCTGAGAAACAGATGCTTGTGTTCTGAAG 300
Db    241  GGAAGAACTTGACTATGGCTCCATGACAGAAAGCTGAGAAACAGATGCTTGTGTTCTGAAG 300

Qy    301  TGAATTGTGCTTCGTGAACCTGAACATCCAAACATCGTTCGTACTATGATCGGATATTG 360
Db    301  TGAATTGTGCTTCGTGAACCTGAACATCCAAACATCGTTCGTACTATGATCGGATATTG 360

Qy    361  ACCGGACCAATACAACTGTACATTGTAATGGAATATTGTGAAGGAGGGGATCTGGCTA 420
Db    361  ACCGGACCAATACAACTGTACATTGTAATGGAATATTGTGAAGGAGGGGATCTGGCTA 420

Qy    421  GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTCTTCGAG 480
Db    421  GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTCTTCGAG 480

Qy    481  TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA 540
Db    481  TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA 540

Qy    541  CCGTATTGCATCGGGATCTTAAACAGCCAAATGTTTTCTGGATGGCAAGCAAAACGTCA 600
Db    541  CCGTATTGCATCGGGATCTTAAACAGCCAAATGTTTTCTGGATGGCAAGCAAAACGTCA 600

Qy    601  AGCTTGGAGACTTTGGGCTAGCTAGAATATTAACCATGACACAGAGTTTGGCAAAACAT 660
Db    601  AGCTTGGAGACTTTGGGCTAGCTAGAATATTAACCATGACACAGAGTTTGGCAAAACAT 660

Qy    661  TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACATGAGA 720
Db    661  TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACATGAGA 720

Qy    721  AATCAGATATCTGGTCATTGGGCTGCTGTGTATGAGTTATGTGCATTAAATGCTCCAT 780

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Db	721		AATCAGATATCTGGTCATTGGGCTGCTGCTGTATGAGTTATGTGCATTAACTGCTCCAT	780
Qy	781		TTACAGCTTTTAGCCAGAAAGAACTCGCTGGGAAAAACAGAGAAGGCAAAATTCAGGCGAA	840
Db	781		TTACAGCTTTTAGCCAGAAAGAACTCGCTGGGAAAAACAGAGAAGGCAAAATTCAGGCGAA	840
Qy	841		TTCCATACCGTTACTCTGATGAATGAATGAATATTATACAGGATGTTAACTTAAAG	900
Db	841		TTCCATACCGTTACTCTGATGAATGAATGAATATTATACAGGATGTTAACTTAAAG	900
Qy	901		ATTACCATCGACCTTCTGTTGAAGAAATCTTGAAGACCCCTTTAATAGCAGATTGGTGT	960
Db	901		ATTACCATCGACCTTCTGTTGAAGAAATCTTGAAGACCCCTTTAATAGCAGATTGGTGT	960
Qy	961		CAGACGAGCAAGAGAAATCTTGAGAGAAAGAGGCCACAATTAGGAGAGCCAGAAAAAT	1020
Db	961		CAGACGAGCAAGAGAAATCTTGAGAGAAAGAGGCCACAATTAGGAGAGCCAGAAAAAT	1020
Qy	1021		CCGAGGATTCCAGCCCTGTATTGAGTGACCTGAAGCTGAAGCAATTCAGTTACAGGAGC	1080
Db	1021		CCGAGGATTCCAGCCCTGTATTGAGTGAGCTGAAGCTGAAGCAATTCAGTTACAGGAGC	1080
Qy	1081		GAGAGCGAGCTCTCAAAGCAAGAGAAAGAAATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Db	1081		GAGAGCGAGCTCTCAAAGCAAGAGAAAGAAATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Qy	1141		TTGTTGAGAGACTAGCAGAGGACAACTGGCTAGAGCAGAAAACTGTTGAAGAACTACA	1200
Db	1141		TTGTTGAGAGACTAGCAGAGGACAACTGGCTAGAGCAGAAAACTGTTGAAGAACTACA	1200
Qy	1201		GCTTCTTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTCTTAACTTTC	1260
Db	1201		GCTTCTTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTCTTAACTTTC	1260
Qy	1261		CATCTCTCAGTAATTAAAGAAAGTTCAATTCAGTGGGGAAGTAAGAGAACATCATGA	1320
Db	1261		CATCTCTCAGTAATTAAAGAAAGTTCAATTCAGTGGGGAAGTAAGAGAACATCATGA	1320
Qy	1321		GGAGTGAGAAATTCTGAGATCAGCTCAGATCACTAAGTCCAAAGTCAAGGACCTGAAGAAA	1380
Db	1321		GGAGTGAGAAATTCTGAGATCAGCTCAGATCACTAAGTCCAAAGTCAAGGACCTGAAGAAA	1380
Qy	1381		GGCTTCACGCTGCCAGCTGCGGGCTCAAGCCCTGTGAGATATTGAGAAAAATACCAC	1440
Db	1381		GGCTTCACGCTGCCAGCTGCGGGCTCAAGCCCTGTGAGATATTGAGAAAAATACCAC	1440
Qy	1441		TGAAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGACACAGAGCTGTGTAC	1500
Db	1441		TGAAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGACACAGAGCTGTGTAC	1500
Qy	1501		AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAACTACT	1560
Db	1501		AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAACTACT	1560
Qy	1561		GGCCCCATGAGCCATGCGCTTCTGTATAGTACACATGATATTCGGAATTGGTTTACTG	1620
Db	1561		GGCCCCATGAGCCATGCGCTTCTGTATAGTACACATGATATTCGGAATTGGTTTACTG	1620
Qy	1621		TTCTTCAGCAACTATTGTACAAAATGTTACATTTAAATTTTCTTTCTCTTTTAAAGAC	1680
Db	1621		TTCTTCAGCAACTATTGTACAAAATGTTACATTTAAATTTTCTTTCTCTTTTAAAGAC	1680
Qy	1681		ATATTATAAAAAAGAACTCTTCTGGTGGGCTTTAATCCTGTGTGTGATTACTAGTAG	1740
Db	1681		ATATTATAAAAAAGAACTCTTCTGGTGGGCTTTAATCCTGTGTGTGATTACTAGTAG	1740
Qy	1741		GAACATGAGATGTGACATCTTAAATCTTGGGAGAAAAATAATATTAGAAAAAAATATT	1800
Db	1741		GAACATGAGATGTGACATCTTAAATCTTGGGAGAAAAATAATATTAGAAAAAAATATT	1800
Qy	1801		TATGAGGAAGAGTAGCACTCACTGAATAGTTTAAATGACTGAGTGGTATGCTTACAA	1860
Db	1801		TATGAGGAAGAGTAGCACTCACTGAATAGTTTAAATGACTGAGTGGTATGCTTACAA	1860
Qy	1861		TGTCATGCTAGATTTAAATTTTAAAGTCTGAGATTTTAAATGTTTTGAGCTTAGAAAA	1920

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Db      1861  |||||
1861  TGTGATGCTAGATTTAAATTTTAAAGCTGAGATTTTAAATGTTTTGAGCTTAGAAAAC 1920

Qy      1921  CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAAATATTCATTGC 1980
1921  |||||

Db      1921  CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAAATATTCATTGC 1980

Qy      1981  TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCCTTT 2040
1981  |||||

Db      1981  TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCCTTT 2040

Qy      2041  TTTTCTCTGTTTAACAGAAATATGAGCTGTCTGTGATTTACCTACTTCTTCCACTAAA 2100
2041  |||||

Db      2041  TTTTCTCTGTTTAACAGAAATATGAGCTGTCTGTGATTTACCTACTTCTTCCACTAAA 2100

Qy      2101  TAAAAGAAATCTTCAGTTA 2119
2101  |||||

Db      2101  TAAAAGAAATCTTCAGTTA 2119
<!--EndFragment-->

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